RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: _____ 10/664, 234 B _____

Source: _____ IFW/6, _____

Date Processed by STIC: _____ 12/15/2006 _____

ENTERED

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        RAW SEQUENCE LISTING                    DATE: 12/15/2006
        PATENT APPLICATION:  US/10/664,234B     TIME: 09:40:11

        Input Set : A:\3240-105.ST25.txt
        Output Set: N:\CRF4\12152006\J664234B.raw
```

```
     3  <110> APPLICANT: Ruan, Yijun
     4        Ng, Patrick
     5        Wei, Chialin
     7  <120> TITLE OF INVENTION: Method for Gene Identification Signature (GIS) Analysis
     9  <130> FILE REFERENCE: 3240-105
C--> 11  <140> CURRENT APPLICATION NUMBER: US/10/664,234B
    12  <141> CURRENT FILING DATE: 2003-09-17
    14  <160> NUMBER OF SEQ ID NOS: 29
    16  <170> SOFTWARE: PatentIn version 3.3
    18  <210> SEQ ID NO: 1
    19  <211> LENGTH: 33
    20  <212> TYPE: DNA
    21  <213> ORGANISM: Artificial
    23  <220> FEATURE:
    24  <223> OTHER INFORMATION: oligonucleotide primer with homology to bacterial cloning
vector
    27  <220> FEATURE:
    28  <221> NAME/KEY: misc_feature
    29  <222> LOCATION: (1)..(33)
    30  <223> OTHER INFORMATION: n is a,c,g, or t
    32  <220> FEATURE:
    33  <221> NAME/KEY: misc_feature
    34  <222> LOCATION: (1)..(33)
    35  <223> OTHER INFORMATION: v is a,c,g
    37  <400> SEQUENCE: 1
W--> 38  gagctccttc tggagttttt tttttttttt tvn                           33
    41  <210> SEQ ID NO: 2
    42  <211> LENGTH: 30
    43  <212> TYPE: DNA
    44  <213> ORGANISM: Artificial
    46  <220> FEATURE:
    47  <223> OTHER INFORMATION: oligonucleotide primer with homology to bacterial cloning
vector
    50  <220> FEATURE:
    51  <221> NAME/KEY: misc_feature
    52  <222> LOCATION: (1)..(30)
    53  <223> OTHER INFORMATION: n is a,t,c or g
    55  <400> SEQUENCE: 2
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    60  <211> LENGTH: 20
    61  <212> TYPE: DNA
    62  <213> ORGANISM: Artificial
    64  <220> FEATURE:
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65 <223> OTHER INFORMATION: oligonucleotide primer with homology to bacterial cloning vector

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Input Set : A:\3240-105.ST25.txt
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 67 <400> SEQUENCE: 3
 68 gtcggatcca agcggccgcg                                          20
 71 <210> SEQ ID NO: 4
 72 <211> LENGTH: 30
 73 <212> TYPE: DNA
 74 <213> ORGANISM: Artificial
 76 <220> FEATURE:
 77 <223> OTHER INFORMATION: oligonucleotide primer with homology to bacterial cloning
vector
 80 <220> FEATURE:
 81 <221> NAME/KEY: misc_feature
 82 <222> LOCATION: (1)..(30)
 83 <223> OTHER INFORMATION: n is a,t,c or g
 85 <400> SEQUENCE: 4
W--> 86 aattcgcggc cgcttggatc cgacgnnnnn                            30
 89 <210> SEQ ID NO: 5
 90 <211> LENGTH: 19
 91 <212> TYPE: DNA
 92 <213> ORGANISM: Artificial
 94 <220> FEATURE:
 95 <223> OTHER INFORMATION: oligonucleotide primer with homology to bacterial cloning
vector
 97 <400> SEQUENCE: 5
 98 tcgacccagg atccaactt                                           19
101 <210> SEQ ID NO: 6
102 <211> LENGTH: 13
103 <212> TYPE: DNA
104 <213> ORGANISM: Artificial
106 <220> FEATURE:
107 <223> OTHER INFORMATION: oligonucleotide primer with homology to bacterial cloning
vector
110 <220> FEATURE:
111 <221> NAME/KEY: misc_feature
112 <222> LOCATION: (1)..(13)
113 <223> OTHER INFORMATION: phosporylation
115 <400> SEQUENCE: 6
116 gttggatcct ggg                                                 13
119 <210> SEQ ID NO: 7
120 <211> LENGTH: 17
121 <212> TYPE: DNA
122 <213> ORGANISM: Artificial
124 <220> FEATURE:
125 <223> OTHER INFORMATION: oligonucleotide primer with homology to bacterial cloning
vector
127 <400> SEQUENCE: 7
128 gtaaaacgac ggccagt                                             17
131 <210> SEQ ID NO: 8
132 <211> LENGTH: 19
133 <212> TYPE: DNA
134 <213> ORGANISM: Artificial
136 <220> FEATURE:
137 <223> OTHER INFORMATION: oligonucleotide primer with homology to bacterial cloning
vector
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139 <400> SEQUENCE: 8

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    146 <213> ORGANISM: Artificial
    148 <220> FEATURE:
    149 <223> OTHER INFORMATION: oligonucleotide primer with homology to bacterial cloning
vector
    151 <400> SEQUENCE: 9
    152 taatacgact cactataggg                                             20
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    156 <211> LENGTH: 22
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    158 <213> ORGANISM: Artificial
    160 <220> FEATURE:
    161 <223> OTHER INFORMATION: oligonucleotide primer with homology to bacterial cloning
vector
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    167 <210> SEQ ID NO: 11
    168 <211> LENGTH: 23
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    170 <213> ORGANISM: Artificial
    172 <220> FEATURE:
    173 <223> OTHER INFORMATION: oligonucleotide primer with homology to bacterial cloning
vector
    175 <400> SEQUENCE: 11
    176 agcggataac aatttcacac agg                                         23
    179 <210> SEQ ID NO: 12
    180 <211> LENGTH: 48
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    182 <213> ORGANISM: Artificial
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    188 <220> FEATURE:
    189 <221> NAME/KEY: misc_feature
    190 <222> LOCATION: (1)..(48)
    191 <223> OTHER INFORMATION: n is a,t,c or g
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    198 <211> LENGTH: 48
    199 <212> TYPE: DNA
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    202 <220> FEATURE:
    203 <223> OTHER INFORMATION: Oligionucleotide with homolgy to a bacteria cloning vector
    206 <220> FEATURE:
    207 <221> NAME/KEY: misc_feature
    208 <222> LOCATION: (1)..(48)
    209 <223> OTHER INFORMATION: n is a,t,c or g
    211 <400> SEQUENCE: 13
W--> 212 gatccaactt nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnngtcg              48
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RAW SEQUENCE LISTING   DATE: 12/15/2006
PATENT APPLICATION: US/10/664,234B   TIME: 09:40:11

Input Set : A:\3240-105.ST25.txt
Output Set: N:\CRF4\12152006\J664234B.raw

215 <210> SEQ ID NO: 14
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218 <213> ORGANISM: Artificial
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221 <223> OTHER INFORMATION: Oligionucleotide primer with homolgy to a bacteria cloning
vector
224 <220> FEATURE:
225 <221> NAME/KEY: misc_feature
226 <222> LOCATION: (1)..(29)
227 <223> OTHER INFORMATION: phosphorylation
229 <400> SEQUENCE: 14
230 cgctctcctg taccgaccct gccgcttac   29
233 <210> SEQ ID NO: 15
234 <211> LENGTH: 29
235 <212> TYPE: DNA
236 <213> ORGANISM: Artificial
238 <220> FEATURE:
239 <223> OTHER INFORMATION: Oligionucleotide primer with homolgy to a bacteria cloning
vector
242 <220> FEATURE:
243 <221> NAME/KEY: misc_feature
244 <222> LOCATION: (1)..(29)
245 <223> OTHER INFORMATION: phosphorylation
247 <400> SEQUENCE: 15
248 aactatcgtc ttgagaccaa cccggtaag   29
251 <210> SEQ ID NO: 16
252 <211> LENGTH: 24
253 <212> TYPE: DNA
254 <213> ORGANISM: Artificial
256 <220> FEATURE:
257 <223> OTHER INFORMATION: Oligionucleotide adapter with homolgy to a bacteria cloning
258   vector
260 <400> SEQUENCE: 16
261 aattctcgag cggccgcgat atcg   24
264 <210> SEQ ID NO: 17
265 <211> LENGTH: 24
266 <212> TYPE: DNA
267 <213> ORGANISM: Artificial
269 <220> FEATURE:
270 <223> OTHER INFORMATION: Oligionucleotide adapter with homolgy to a bacteria cloning
271   vector
274 <220> FEATURE:
275 <221> NAME/KEY: misc_feature
276 <222> LOCATION: (1)..(24)
277 <223> OTHER INFORMATION: phosphorylation
279 <400> SEQUENCE: 17
280 aattcgatat cgcggccgct cgag   24
283 <210> SEQ ID NO: 18
284 <211> LENGTH: 3404
285 <212> TYPE: DNA

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286 <213> ORGANISM: Artificial
288 <220> FEATURE:
289 <223> OTHER INFORMATION: bacterial cloning vector
291 <400> SEQUENCE: 18
292 gggcgaattc tcgagcggcc gcggatccga cgagagcgcc tgcgtacggc tcgccgcggt      60
294 ggctggcgct acttcggagg agcccgacgc ggcgcggtcg tttttataca ttcccgcgcg     120
296 gaggcaacgg aagggcgggg cgcctcgtga ttaggccgcg gaggtcacag gctctgttgt     180
298 catgaaggtg aaaattaaat gttggaatgg tgtggccact tggctctggg tagccaatga     240
300 tgagaactgc ggcatctgca ggatggcgtt taatggctgc tgtccagact gtaaggtgcc     300
302 tggtgatgac tgccccctcg tgtgggggaca gtgctcccac tgcttccaca tgcactgcat     360
304 cctcaagtgg ctgaatgcgc agcaggtgca gcagcactgc cccatgtgtc gccaggagtg     420
306 gaagttcaaa gagtgaagcc cgtgccgtgc cacttccctc tcctgtgctg tgccaggctc     480
308 agccccttcc ctccctcccc tccccagat acagcacccc aagtcccctc cacacagcac     540
310 agtggtgccc agagatctcg gtctgtgccg gggacaagga tgctttctgt ttggctggga     600
312 caaggttgaa aggagctttg ctgactgttt tgttttccca tcacattgac actttattca     660
314 ataagtaaaa ctcattacag ttccaagtcg gatcctgggt cgacctgcag gcatgcaagc     720
316 ttgagtattc tatagtgtca cctaaatagc ttggcgtaat catggtcata gctgtttcct     780
318 gtgtgaaatt gttatccgct cacaattcca cacaacatac gagccggaag cataaagtgt     840
320 aaagcctggg gtgcctaatg agtgagctaa ctcacattaa ttgcgttgcg ctcactgccc     900
322 gctttccagt cgggaaacct gtcgtgccag ctgcattaat gaatcggcca acgcgcgggg     960
324 agaggcggtt tgcgtattgg gcgctcttcc gcttcctcgc tcactgactc gctgcgctcg    1020
326 gtcgttcggc tgcggcgagc ggtatcagct cactcaaagg cggtaatacg gttatccaca    1080
328 gaatcagggg ataacgcagg aaagaacatg tgagcaaaag gccagcaaaa ggccaggaac    1140
330 cgtaaaaagg ccgcgttgct ggcgtttttc gataggctcc gcccccctga cgagcatcac    1200
332 aaaaatcgac gctcaagtca gaggtggcga aacccgacag gactataaag ataccaggcg    1260
334 tttcccctg gaagctccct cgtgcgctct cctgtaccga ccctgccgct taccggatac    1320
336 ctgtccgcct ttctcccttc gggaagcgtg gcgctttctc atagctcacg ctgtaggtat    1380
338 ctcagttcgg tgtaggtcgt tcgctccaag ctgggctgtg tgcacgaacc ccccgttcag    1440
340 cccgaccgct gcgccttatc cggtaactat cgtcttgaga ccaacccggt aagacacgac    1500
342 ttatcgccac tggcagcagc cactggtaac aggattagca gagcgaggta tgtaggcggt    1560
344 gctacagagt tcttgaagtg gtggcctaac tacggctaca ctagaaggac agtatttggt    1620
346 atctgcgctc tgctgaagcc agttaccttc ggaaaaagag ttggtagctc ttgatccggc    1680
348 aaacaaacca ccgctggtag cggtggtttt tttgtttgca agcagcagat tacgcgcaga    1740
350 aaaaaaggat ctcaagaaga tcctttgatc ttttctacgg ggtctgacgc tcagtggaac    1800
352 gaaaactcac gttaagggat tttggtcatg agattatcaa aaaggatctt cacctagatc    1860
354 cttttaaatt aaaaatgaag ttttaaatca atctaaagta tatatgagta aacttggtct    1920
356 gacagttacc aatgcttaat cagtgaggca cctatctcag cgatctgtct atttcgttca    1980
358 tccatagttg cctgactccc cgtcgtgtag ataactacga tacgggaggg cttaccatct    2040
360 ggccccagtg ctgcaatgat accgcgagac ccacgctcac cggctccaga tttatcagca    2100
362 ataaaccagc cagccggaag ggccgagcgc agaagtggtc ctgcaacttt atccgcctcc    2160
364 atccagtcta ttaattgttg ccgggaagct agagtaagta gttcgccagt taatagtttg    2220
366 cgcaacgttg ttggcattgc tacaggcatc gtggtgtcac gctcgtcgtt ggtatggct    2280
368 tcattcagct ccggttccca acgatcaagg cgagttacat gatccccat gttgtgcaaa    2340
370 aaagcggtta gctccttcgg tcctccgatc gttgtcagaa gtaagttggc cgcagtgtta    2400
372 tcactcatgg ttatggcagc actgcataat tctcttactg tcatgccatc cgtaagatgc    2460
374 ttttctgtga ctggtgagta ctcaaccaag tcattctgag aatagtgtat gcggcgaccg    2520
376 agttgctctt gcccggcgtc aatacgggat aataccgcgc cacatagcag aactttaaaa    2580
378 gtgctcatca ttggaaaacg ttcttcgggg cgaaaactct caaggatctt accgctgttg    2640
380 agatccagtt cgatgtaacc cactcgtgca cccaactgat cttcagcatc ttttactttc    2700
```

RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 12/15/2006
PATENT APPLICATION: US/10/664,234B   TIME: 09:40:12

Input Set : A:\3240-105.ST25.txt
Output Set: N:\CRF4\12152006\J664234B.raw

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 33
Seq#:2; N Pos. 25,26,27,28,29,30
Seq#:4; N Pos. 26,27,28,29,30
Seq#:12; N Pos. 9,10,11,12,13,14,15,16,17,18,19,20,21,22,23,24,25,26,27,28
Seq#:12; N Pos. 29,30,31,32,33,34,35,36,37,38,39,40,41,42
Seq#:13; N Pos. 11,12,13,14,15,16,17,18,19,20,21,22,23,24,25,26,27,28,29,30
Seq#:13; N Pos. 31,32,33,34,35,36,37,38,39,40,41,42,43,44

Invalid <213> Response:
Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,23,24,25,27,28
Seq#:29

VERIFICATION SUMMARY   DATE: 12/15/2006
PATENT APPLICATION: **US/10/664,234B**   TIME: 09:40:12

Input Set : **A:\3240-105.ST25.txt**
Output Set: **N:\CRF4\12152006\J664234B.raw**

```
L:11 M:270 C: Current Application Number differs, Replaced Current Application Number
L:38 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
L:56 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0
L:86 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0
L:194 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0
L:212 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0
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